What is claimed is:

- 1. A mass spectrometry process for analyzing substance mixtures using a triple quadrupole mass spectrometer, said substance mixtures being ionized before the analysis, which comprises the following steps:
- a) selecting a mass/charge quotient (m/z) of an ion formed

 by ionization in a first analytical quadrupole (I) of the

 mass spectrometer,
- b) fragmenting the ion selected under (a) by applying an acceleration voltage in a further following quadrupole
 (II) which is filled with a collision gas and functions as a collision chamber,
 - c) selecting a mass/charge quotient of an ion formed by the fragmentation (b) in a further downstream quadrupole (III), the process steps (a) to (c) being run through at least once, and
- analyzing the mass/charge quotients of all ions present in the substance mixture as a result of the ionization, the quadrupole (II) being filled with collision gas but no acceleration voltage being applied during the analysis;
- and the steps (a) to (c) and step (d) may also be carried out in reverse sequence.
 - 2. The process according to claim 1, wherein the ionization of the substance mixture is upstream of a chromatographic separation.
 - 3. The process according to claim 1 or 2, wherein the chromatographic separation is an HPLC separation.
- 4. The process according to claims 1 to 3, wherein steps (a) to40 (d) are run through at least once within from 0.1 to 10 seconds.

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- 5. The process according to claims 1 to 4, wherein steps (a) to (d) are run through at least once within from 0.2 to 2 seconds.
- 5 6. The process according to claims 1 to 5, wherein the ionization is effected by evaporating the substance mixture and ionizing in the gas phase, by desorbing the substance mixture on a surface or by atomizing the substance mixture in an electrical field.

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- 7. The process according to claims 1 to 6, wherein the ionization is effected by atomizing the substance mixture in an electrical field.
- 15 8. The process according to claims 1 to 7, wherein analysis is effected in step (a) between 1 and 100 mass/charge quotients of different ions formed by ionization and selected.
- 9. The process according to claims 1 to 8, wherein the substance mixture is of biological or chemical origin.
 - 10. The process according to claims 1 to 9, wherein the substance mixtures are derivatized before the analysis or before the chromatographic separation according to claim 2 or 3.

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- 11. The process according to claims 1 to 10, which is carried out manually or automatically.
- 12. The process according to claims 1 to 11, which is used in a high-throughput screening.
- 13. The process according to claims 1 to 12, wherein the fragment ion analyzed in step (c) and the (m/z) quotients, analyzed in step (d), of all ions present in the substance mixture or the fragment ion analyzed in step (c) or the (m/z) quotients, analyzed in step (d), of all ions present in the substance mixture are quantified.

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Mass spectrometry process for analyzing substance mixtures

Abstract

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A mass spectrometry process for analyzing substance mixtures using a triple quadrupole mass spectrometer, said substance mixtures being ionized before the analysis, which comprises the following steps:

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- a) selecting a mass/charge quotient (m/z) of an ion formed by ionization in a first analytical quadrupole (I) of the mass spectrometer,
- 15 b) fragmenting the ion selected under (a) by applying an acceleration voltage in a further following quadrupole (II) which is filled with a collision gas and functions as a collision chamber,
- 20 c) selecting a mass/charge quotient of an ion formed by the fragmentation (b) in a further downstream quadrupole (III), the process steps (a) to (c) being run through at least once, and
- 25 d) analyzing the mass/charge quotients of all ions present in the substance mixture as a result of the ionization, the quadrupole (II) being filled with collision gas but no acceleration voltage being applied during the analysis;
- 30 and the steps (a) to (c) and step (d) may also be carried out in reverse sequence.

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